

IN THE CLAIMS:

This listing of claims will replace all prior versions, and listings, of claims in the application:

1.-57. Cancelled

58. (Previously Presented) An isolated polypeptide comprising the amino acid sequence in SEQ ID NO:13.

59. (Previously Presented) An isolated polypeptide comprising the amino acid sequence encoded by the nucleotide sequence of SEQ ID NO:12.

60. (Currently Amended) An isolated polypeptide ~~comprising amino acid residues 125 to 158 of SEQ ID NO:13~~ which is encoded by a nucleic acid molecule which hybridizes to the complement of ~~[[a]] the nucleic acid molecule that encodes a polypeptide comprising the amino acid sequence of SEQ ID NO:13 in~~ SEQ ID NO:12 under hybridization conditions comprising incubation in 6XSSC at 45°C, followed by one or more washes in 0.2xSSC/0.1% SDS at 50-65°C, wherein the polypeptide comprises amino acid residues 150 to 158 of SEQ ID NO:13.

61. (Canceled)

62. (Currently Amended) ~~An isolated polypeptide comprising amino acid residues 125 to 158 of SEQ ID NO:13 which is encoded by a nucleic acid molecule which hybridizes to the complement of a nucleic acid molecule that encodes a polypeptide comprising the amino acid sequence of SEQ ID NO:13 under~~ The isolated polypeptide of claim 60, wherein the hybridization conditions comprising comprise incubation in 6XSSC at 45°C followed by one or more washes in 0.1xSSC/0.2% SDS at 68°C.

63. (Canceled)

64. (Previously Presented) An isolated polypeptide comprising at least amino acid residues:

- (a) 125 to 158 of SEQ ID NO:13;
- (b) 100 to 158 of SEQ ID NO:13;
- (c) 75 to 158 of SEQ ID NO:13;
- (d) 50 to 158 of SEQ ID NO:13; or

(e) 25 to 158 of SEQ ID NO:13.

65. (Previously Presented) The isolated polypeptide of Claim 58, 59, 60, 62 or 64 further comprising a heterologous polypeptide.

66. (Previously Presented) The isolated polypeptide of Claim 65, wherein the heterologous polypeptide is an Ig polypeptide.

67. (New) A method for identifying a compound which binds to a polypeptide selected from the group consisting of:

i) an isolated polypeptide which is encoded by a nucleic acid molecule which hybridizes to the complement of the nucleic acid molecule in SEQ ID NO:12 under hybridization conditions comprising incubation in 6XSSC at 45°C, followed by one or more washes in 0.2xSSC/0.1% SDS at 50-65°C, wherein the polypeptide comprises amino acid residues 150 to 158 of SEQ ID NO:13;

ii) an isolated polypeptide comprising at least amino acid residues 125 to 158 of SEQ ID NO:13; and

iii) a polypeptide of i) or ii), further comprising heterologous amino acid sequences,

comprising the steps of:

a) contacting a reaction mixture comprising the polypeptide with a test compound; and

b) determining whether the polypeptide binds to the test compound.

68. (New) The method of claim 67, wherein the binding of the test compound to the polypeptide is detected by a method selected from the group consisting of:

a) detection of binding by direct detecting of test compound/polypeptide binding;

b) detection of binding using a competition binding assay; and

c) detection of binding using a two-hybrid assay.

69. (New) The method of claim 67, wherein the polypeptide comprises the amino acid sequence of SEQ ID NO:13.

70. (New) The method of claim 67, wherein the polypeptide is immobilized on a solid surface.

71. (New) The method of claim 67, wherein the test compound is directly or indirectly labeled.